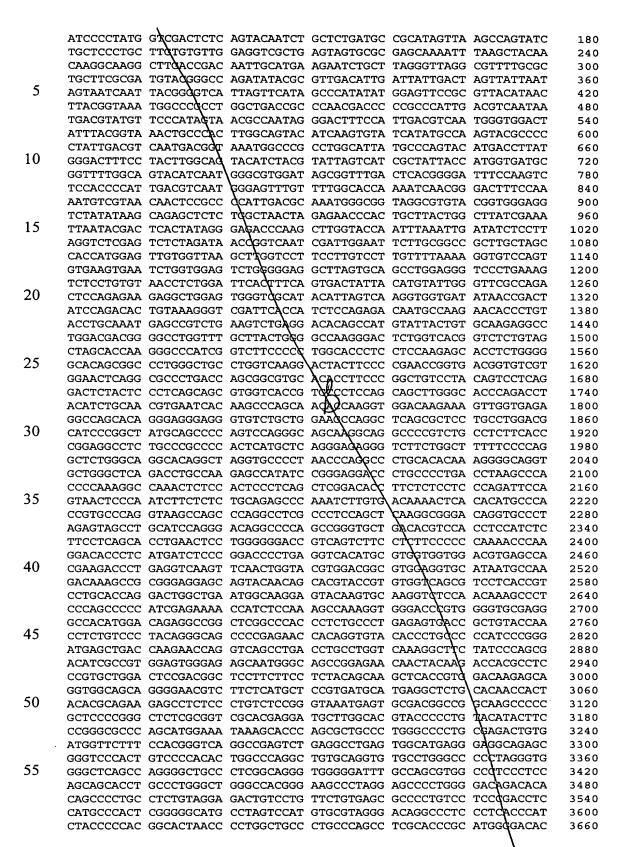
SEQUENCE LISTING

5	(1) GENERAL INFORMATION
3	(i) APPLICANT: Bristol-Myers Squibb Co.
10	(ii) TITLE OF THE INVENTION: A METHOD FOR INHIBITING IMMUNOGLOBULIN-INDUCED TOXICITY FROM THE USE OF IMMUNOGLOBULINS IN THERAPY AND IN VIVO DIAGNOSIS
15	(iii) NUMBER OF SEQUENCES: 13(iv) CORRESPONDENCE ADDRESS:(A) ADDRESSEE: Merchant & Gould(B) STREET: 11150 Santa Monica Blvd., Suite 400
20	(C) CITY: Los Angeles (D) STATE: CA (E) COUNTRY: USA (F) ZIP: 90025
25	(v) COMPUTER READABLE FORM: (A) MEDIUM TYPE: Diskette (B) COMPUTER: IBM Compatible (C) OPERATING SYSTEM: DOS (D) SOFTWARE: FastSEQ Version 2.0
30 000	(vi) CURRENT APPLICATION DATA: (A) APPLICATION NUMBER: PCT/US93/ (B) FILING DATE: 01-AUG-1997 (C) CLASSIFICATION:
35	(vii) PRIOR APPLICATION DATA: (A) APPLICATION NUMBER: 60/023,033 (B) FILING DATE: 02-AUG-1996
40	(viii) ATTORNEY/AGENT INFORMATION: (A) NAME: Adriano, Sarah B (B) REGISTRATION NUMBER: 34,470 (C) REFERENCE/DOCKET NUMBER: 30436.43WOU1
45	(ix) TELECOMMUNICATION INFORMATION: (A) TELEPHONE: 310-445-1140 (B) TELEFAX: 310-445-9031 (C) TELEX:
50	(2) INFORMATION FOR SEQ ID NO:1:
55	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear
	,

	•	
	(ii) MOLECULE TYPE: cDNA	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:	
5	TGGCACCGAA AGCTTTCTGG GGCAGGCCAG GCCTGA	36
	(2) INFORMATION FOR SEQ ID NO:2:	
	(i) SEQUENCE CHARACTERISTICS:	
10	(A) LENGTH: 5V base pairs	
	(B) TYPE: nucleic acid (C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
15	(ii) MOLECULE TYPE: CDNA	
	(vi) SEQUENCE DESCRIPTION, SEQ ID NO.2.	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:	
20	TCCGGACATG TTGGTACCCA CGTGGTGGTC GACGCTGAGC CTGGCTTCGA GCAGACA	57
	(2) INFORMATION FOR SEQ ID NO:3:	
	(i) SEQUENCE CHARACTERISTICS:	
25	(A) LENGTH: 55 base pairs	
25	(B) TYPE: nucleic acid (C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: cDNA	
30	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:	
	GTCGACCACC ACGTGGGTAC CAACATGTCC GGAGCCACAT GGACAGAGGC CGGCT	55
35	(2) INFORMATION FOR SEQ ID NO 4:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 30 base pairs (B) TYPE: nucleic acid	
40	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: cDNA	
45	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:	
	CTGGTTCTTG TTCATCTCCT CTCTAGATGG	30
50	(2) INFORMATION FOR SEQ ID NO:5:	
50	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 36 base pairs	
	(B) TYPE: nucleic acid (C) STRANDEDNESS: single	
55	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: cDNA	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:	
	\cdot	
	51	

	ACCATGGTCG ACCTCAGACC TGCCAAGAGC CATATC	36
5	(2) INFORMATION FOR SEQ ID NO:6:	
3	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH:\\39 base pairs	
	(B) TYPE: nucleic acid	
10	(C) STRANDEDNESS: single (D) TOPOLOGY: linear	
10	(D) TOPOLOGI: tillear	
	(ii) MOLECULE TYPE: CDNA	
1.5	(xi) SEQUENCE DESCRAPTION: SEQ ID NO:6:	
15	CATGGTCACG TGGTGTCC CTGGATGCAG GCTACTCTA	39
	(2) INFORMATION FOR SEQ ID NO:7:	
	(2) INFORMATION FOR SEQ ID NO: /:	
20	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 49 base pairs	
	(B) TYPE: nucleic acid \	
	(C) STRANDEDNESS: single \ (D) TOPOLOGY: linear	
25	(b) TOPOLOGI: Illieal	
	(ii) MOLECULE TYPE: cDNA	
	(xi) SEQUENCE DESCRIPTION: SECOLD NO:7:	
30	CAGGGAGGA GGGTGTCTGC TGGAAGCCAG GCTCAGCT GACCTCAGA	49
	(2) INFORMATION FOR SEQ ID NO:\8:	
35	(i) SEQUENCE CHARACTERISTICS:	
33	(A) LENGTH: 50 base pairs (B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
40		
40	(ii) MOLECULE TYPE: cDNA	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:	
15	GGAAAGAACC ATCACAGTCT CGCAGGGGCC CAGGGCAGCG CTGGGTGCTT	50
45	(2) INFORMATION FOR SEC ID NO. 9.	
	(2) INFORMATION FOR SEQ ID NO:9:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 8691 base pairs	
50	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: cDNA	
55		
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:	
	GACGGATCGG GAGATCTGCT AGGTGACCTG AGGCGCGCCG GCTTCGAATA GCCAGAGTAA	60
	CCTTTTTTT TAATTTTATT TTATTTTATT TTTGAGATGG AGTTTGGCGC CGATCTCCCG	120
		



	\						
	AACCGACTCC	GGGGACATGC	ACTCTCGGGC	CCTGTGGAGG	GACTGGTGCA	GATGCCCACA	3720
	CACACACTCA	&CCCAGACCC	GTTCAACAAA	CCCCGCACTG	AGGTTGGCCG	GCCACACGGC	3780
	CACCACACAC	ACACGTGCAC	GCCTCACACA	CGGAGCCTCA	CCCGGGCGAA	CTGCACAGCA	3840
_			TCGCACACGT				3900
5	CCCCACGCGG	CACCTCAAGG	CCCACGAGCC	TCTCGGCAGC	TTCTCCACAT	GCTGACCTGC	3960
	TCAGACAAAC	CCAGCCTCC	TCTCACAAGG	GTGCCCCTGC	AGCCGCCACA	CACACACAGG	4020
	GGATCACACA	CCACGTCACG	TCCCTGGCCC	TGGCCCACTT	CCCAGTGCCG	CCCTTCCCTG	4080
	CAGGACGGAT	CAGCCT CGAC	TGTGCCTTCT	AGTTGCCAGC	CATCTGTTGT	TTGCCCCTCC	4140
	CCCGTGCCTT	CCTTGACCCT	GGAAGGTGCC	ACTCCCACTG	TCCTTTCCTA	ATAAAATGAG	4200
10	GAAATTGCAT	CGCATTGT/CT	GAGTAGGTGT	CATTCTATTC	TGGGGGGTGG	GGTGGGGCAG	4260
	GACAGCAAGG	GGGAGGATTG	GGAAGACAAT	AGCAGGCATG	CTGGGGATGC	GGTGGGCTCT	4320
	ATGGCTTCTG	AGGCGGAAAd	AACCAGCTGG	GGCTCTAGGG	GGTATCCCCA	CGCGCCCTGT	4380
	AGCGGCGCAT	TAAGCGCGGC	\GGGTGTGGTG	GTTACGCGCA	GCGTGACCGC	TACACTTGCC	4440
	AGCGCCCTAG	CGCCCGCTCC	TTTCGCTTTC	TTCCCTTCCT	TTCTCGCCAC	GTTCGCCGGG	4500
15	CCTCTCAAAA	AAGGGAAAAA	AAGCATGCAT	CTCAATTAGT	CAGCAACCAT	AGTCCCGCCC	4560
	CTAACTCCGC	CCATCCCGCC	CCTAACTCCG	CCCAGTTCCG	CCCATTCTCC	GCCCCATGGC	4620
	TGACTAATTT	TTTTTATTTA	TGQAGAGGCC	GAGGCCGCCT	CGGCCTCTGA	GCTATTCCAG	4680
	AAGTAGTGAG	GAGGCTTTTT	TGGAGGCCTA	${\tt GGCTTTTGCA}$	AAAAGCTTGG	ACAGCTCAGG	4740
	GCTGCGATTT	CGCGCCAAAC	TTGAÇGCAA	TCCTAGCGTG	AAGGCTGGTA	GGATTTTATC	4800
20	CCCGCTGCCA	TCATGGTTCG	ACCATTGAAC	TGCATCGTCG	CCGTGTCCCA	AAATATGGGG	4860
			ACCCTGGCCT				4920
	AGAATGACCA	CAACCTCTTC	AGTGGAAGGT	AAACAGAATC	TGGTGATTAT	GGGTAGGAAA	4980
	ACCTGGTTCT	CCATTCCTGA	GAAGAATÒGA	CCTTTAAAGG	ACAGAATTAA	TATAGTTCTC	5040
2.5			ACCACGAGGA				5100
25			ACCGGAATT	/1			5160
			GGAAGCCATG				5220
	ACAAGGATCA	TGCAGGAATT	TGAAAGTGAC	CGTTTTTCC	CAGAAATTGA	TTTGGGGAAA	5280
			CCCAGGCGTC	\			5340
20			CGAGAAGAAA	\			5400
30			CATTTTTATA	1			5460
			TTCTGTGGTG				5520
			ATAAAATTT				5580
			ATTCCAACCT	\			5640
25			CTGTTTTGCT	1			5700
35			TCTACTCCTC	\			5760
			CTAAGTTTTT	\			5820
			ACCACAAAGG	\ \			5880
			TTTATAAGTA	\			5940
40			CATAGAGTGT		1		6000
40			TGTAAAGGGG		\		6060
			CCATACCACA		\		6120
			CCTGAAACAT		1		6180
			TTACAAATAA		_ ~~\~ ~~ - ~~		6240
45					\	ATCTTATCAT	6300
43			CCTCCAGCGC		\		6360
			TTATAATGGT ACTGCATTCT		1		6420
			GTCGACCTCT		,		6480 6540
			TTATCCGCTC		\		6600
50			TGCCTAATGA		1		6660
50			GGGAAACCTG		١		6720
			GCGTATTGGG		\		6780
			GCGGCGAGCG		1		6840
			TAACGCAGGA		*		6900
55			CGCGTTGCTG		١,		6960
			CTCAAGTCAG				7020
			AAGCTCCCTC			1	7020
			TCTCCCTTCG			1	7140
			GTAGGTCGTT			1	7200
						-7	0 0

30

35



(2) INFORMATION FOR SEQ ID NO:10:

- (i) SEQUENCE CHARACTERISTICS
 - (A) LENGTH: 8327 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

				١ .			
	GACGGATCGG	GAGATCTGCT	AGGTGACCTG	AGGCGCGCC	GCTTCGAATA	GCCAGAGTAA	60
40	CCTTTTTTTT	TAATTTTATT	TTATTTTATT	TTTGAGATGG	AGTTTGGCGC	CGATCTCCCG	120
	ATCCCCTATG	GTCGACTCTC	AGTACAATCT	GCTCTGATGC	CGCATAGTTA	AGCCAGTATC	180
	TGCTCCCTGC	TTGTGTGTTG	GAGGTCGCTG	AGTAGTGCGC	GAGCAAAATT	TAAGCTACAA	240
	CAAGGCAAGG	CTTGACCGAC	AATTGCATGA	AGAATCTGCT	TAGGGTTAGG	CGTTTTGCGC	300
	TGCTTCGCGA	TGTACGGGCC	AGATATACGC	GTTGACATTG	ATTATTGACT	AGTTATTAAT	360
45	AGTAATCAAT	TACGGGGTCA	TTAGTTCATA	GCCCATATAT	GGAGTTCCGC	GTTACATAAC	420
	TTACGGTAAA	TGGCCCGCCT	GGCTGACCGC	CCAACGACCC	CCGCCATTG	ACGTCAATAA	480
	TGACGTATGT	TCCCATAGTA	ACGCCAATAG	${\tt GGACTTTCCA}$	TTGAÇGTCAA	TGGGTGGACT	540
	ATTTACGGTA	AACTGCCCAC	TTGGCAGTAC	ATCAAGTGTA	TCATATGCCA	AGTACGCCCC	600
	CTATTGACGT	CAATGACGGT	AAATGGCCCG	CCTGGCATTA	TGCCCAGTAC	ATGACCTTAT	660
50	GGGACTTTCC	TACTTGGCAG	TACATCTACG	TATTAGTCAT	CGCTATTACC	ATGGTGATGC	720
	GGTTTTGGCA	GTACATCAAT	GGGCGTGGAT	${\tt AGCGGTTTGA}$	CTCACGGGGA	TTTCCAAGTC	780
	TCCACCCCAT	TGACGTCAAT	GGGAGTTTGT	TTTGGCACCA	AAATCAAdGG	GACTTTCCAA	840
	AATGTCGTAA	CAACTCCGCC	CCATTGACGC	AAATGGGCGG	TAGGCGTGTA	CGGTGGGAGG	900
	TCTATATAAG	CAGAGCTCTC	TGGCTAACTA	GAGAACCCAC	TGCTTACTG	CTTATCGAAA	960
55	TTAATACGAC	TCACTATAGG	GAGACCCAAG	CTTGGTACCA	PTTAAATTTA	ATATCTCCTT	1020
	AGGTCTCGAG	TCTCTAGATA	ACCGGTCAAT	CGATTGGAAT	TCTTGCGGCC	GCTTGCTAGC	1080
	CACCATGGAG	TTGTGGTTAA	GCTTGGTCCT	TCCTTGTCCT	TGTTTTAAAA	\GGTGTCCAGT	1140
	GTGAAGTGAA	TCTGGTGGAG	TCTGGGGGAG	GCTTAGTGCA	GCCTGGAGGG	TCCCTGAAAG	1200
	TCTCCTGTGT	AACCTCTGGA	TTCACTTTCA	GTGACTATTA	CATGTATTGG	GTTCGCCAGA	1260

	λ						
	CTCCAGAGAA	GAGGCTGGAG	TGGGTCGCAT	ACATTAGTCA	AGGTGGTGAT	ATAACCGACT	1320
	ATCCAGACAC	TGTAAAGGGT	CGATTCACCA	TCTCCAGAGA	CAATGCCAAG	AACACCCTGT	1380
	ACCTGCAAAT	GAGCCGTCTG	AAGTCTGAGG	ACACAGCCAT	GTATTACTGT	GCAAGAGGCC	1440
_	TGGACGACGG	GCCTGGTTT	GCTTACTGGG	GCCAAGGGAC	TCTGGTCACG	GTCTCTGTAG	1500
5	CTAGCACCAA	GGGCCCATCG	GTCTTCCCCC	TGGCACCCTC	CTCCAAGAGC	ACCTCTGGGG	1560
	GCACAGCGGC	CCTGGGCTGC	CTGGTCAAGG	ACTACTTCCC	CGAACCGGTG	ACGGTGTCGT	1620
	GGAACTCAGG	CGCQCTGACC	AGCGGCGTGC	ACACCTTCCC	GGCTGTCCTA	CAGTCCTCAG	1680
	GACTCTACTC	CCTCAGCAGC	GTGGTCACCG	TGCCCTCCAG	CAGCTTGGGC	ACCCAGACCT	1740
	ACATCTGCAA	CGTGAATCAC	AAGCCCAGCA	ACACCAAGGT	GGACAAGAAA	GTTGGTGAGA	1800
10	GGCCAGCACA	GGGAGGGAGG	GTGTCTGCTG	GAAGCCAGGC	TCAGCGCTCC	TGCCTGGACG	1860
	CATCCCGGCT	ATGCAGCCC	AGTCCAGGGC	AGCAAGGCAG	GCCCCGTCTG	CCTCTTCACC	1920
	CGGAGGCCTC	recccecqcc	ACTCATGCTC	AGGGAGAGGG	TCTTCTGGCT	TTTTCCCCAG	1980
	GCTCTGGGCA	GGCACAGGÒT	AGGTGCCCCT	AACCCAGGCC	CTGCACACAA	AGGGGCAGGT	2040
1.5	GCTGGGCTCA	GACCTGCCAA	GAGCCATATC	CGGGAGGACC	CTGCCCCTGA	CCTAAGCCCA	2100
15	CCCCAAAGGC	CAAACTCTCC\	ACTCCCTCAG	CTCGGACACC	TTCTCTCCTC	CCAGATTCCA	2160
		ATCTTCTCTC	1				2220
		GTAAGCCAGC	\				2280
		GCATCCAGGG	1				2340
20		CTCGGCCCAC					2400
20		CCCCGAGAAC	1				2460
		GTCAGCCTGA	\				2520
		AGCAATGGGC					2580
		TCCTTCTTCC					2640
25		TTCTCATGCT	\				2700
25		CTGTCTCCGG	١ ـ				2760
		CGCACGAGGA TAAAGCACCC					2820
							2880
		GGCCGAGTCT	• \				2940
30		TGGCCCAGGC	· ·	\			3000
30		CTCGGCAGGG GGGCCACGGG		\			3060
		GACTGTCCTG		\			3120 3180
		CCTAGTCCAT		1			3240
		CCTGGCTGCC		\			3300
35		ACTCTCGGGC		\			3360
55		GTTCAACAAA		1			3420
		GCCTCACACA		\			3480
		TCGCACACGT		\			3540
		CCCACGAGCC		\			3600
40		TCTCACAAGG					3660
		TCCCTGGCCC		`			3720
	CAGCCTCGAC	TGTGCCTTCT	AGTTGCCAGC	CATCTGTTGT	TTGCCCCTCC	CCCGTGCCTT	3780
	CCTTGACCCT	GGAAGGTGCC	ACTCCCACTG	TCCTTTCCTA	\ATAAAATGAG	GAAATTGCAT	3840
	CGCATTGTCT	GAGTAGGTGT	CATTCTATTC	TGGGGGGTGG	GTGGGGCAG	GACAGCAAGG	3900
45	GGGAGGATTG	GGAAGACAAT	AGCAGGCATG	CTGGGGATGC	GGTGGGCTCT	ATGGCTTCTG	3960
	AGGCGGAAAG	AACCAGCTGG	GGCTCTAGGG	GGTATCCCCA	сфсесстет	AGCGGCGCAT	4020
	TAAGCGCGGC	GGGTGTGGTG	GTTACGCGCA	GCGTGACCGC	TACACTTGCC	AGCGCCCTAG	4080
	CGCCCGCTCC	TTTCGCTTTC	TTCCCTTCCT	TTCTCGCCAC	GTT¢GCCGGG	CCTCTCAAAA	4140
	AAGGGAAAAA				\ \		4200
50	CCATCCCGCC	CCTAACTCCG	CCCAGTTCCG	CCCATTCTCC	GCCCCATGGC	TGACTAATTT	4260
	TTTTTATTTA	TGCAGAGGCC	GAGGCCGCCT	CGGCCTCTGA	GCTATT/CCAG	AAGTAGTGAG	4320
		TGGAGGCCTA					4380
		TTGACGGCAA			١		4440
<i></i>		ACCATTGAAC			\		4500
55		ACCCTGGCCT			1		4560
		AGTGGAAGGT					4620
		GAAGAATCGA					4680
		ACCACGAGGA					4740
	TTATTGAACA	ACCGGAATTG	GCAAGTAAAG	TAGACATGGT	TTGGATAGTC	GAGGCAGTT	4800

	\						
	\	-			ACTCTTTGTG		4860
	\				TTTGGGGAAA		4920
	TCCCAGAATA	CCCAGGCGTC	CTCTCTGAGG	TCCAGGAGGA	AAAAGGCATC	AAGTATAAGT	4980
_	\	CGAGAAGAAA	GACTAACAGG	AAGATGCTTT	CAAGTTCTCT	GCTCCCCTCC	5040
5	\	CATTTTTATA	AGACCATGGG	ACTTTTGCTG	GCTTTAGATC	TCTTTGTGAA	5100
	GGAACCTTAC				CCTACAGAGA		5160
	TAAGGTAAAT	\			CTACTGATTC		5220
	TGTATTTTAG	ATTCCAACCT	ATGGAACTGA	TGAATGGGAG	CAGTGGTGGA	ATGCCTTTAA	5280
4.0	TGAGGAAAAC	CTGTTTTGCT	CAGAAGAAAT	GCCATCTAGT	GATGATGAGG	CTACTGCTGA	5340
10	CTCTCAACAT	TCTACTCCTC	CAAAAAAGAA	GAGAAAGGTA	GAAGACCCCA	AGGACTTTCC	5400
	TTCAGAATTG	CTAAGTTTTT	TGAGTCATGC	TGTGTTTAGT	AATAGAACTC	TTGCTTGCTT	5460
		`			AAGAAAATTA		5520
	TTCTGTAACC	TTTATAAGTA	GGCATAACAG	TTATAATCAT	AACATACTGT	TTTTTCTTAC	5580
1.5		\			CAAAAATTGT		5640
15		,	١.		TATAGTGCCT		5700
			1		TTTAAAAAAC		5760
	TCCCCCTGAA	CCTGAAACAT	AAAATGAATG	CAATTGTTGT	TGTTAACTTG	TTTATTGCAG	5820
			`		CACAAATAAA		5880
•	CACTGCATTC	TAGTTGTGGT	TTGTCCAAAC	TCATCAATGT	ATCTTATCAT	GTCTGGATCG	5940
20	GCTGGATGAT	CCTCCAGCGC	GGGGATCTCA	TGCTGGAGTT	CTTCGCCCAC	CCCAACTTGT	6000
			`		CACAAATTTC		6060
			`		CATCAATGTA		6120
			\		ATGGTCATAG		6180
0.5			`		AGCCGGAAGC		6240
25				17	TGCGTTGCGC		6300
					AATCGGCCAA		6360
				`	CACTGACTCG		6420
				1	GGTAATACGG		6480
20				`	CCAGCAAAAG		6540
30				`	CCCCCTGAC		6600
				` `	ACTATAAAGA		6660
				1	CCTGCCGCTT		6720
				`	ATGCTCACGC		6780
2.5				`	GCACGAACCC		6840
35				\	CAACCCGGTA		6900
					AGCGAGGTAT		6960
		CTTGAAGTGG			'TAGAAGGACA		7020
					TGGTAGCTCT		7080
40					GCAGCAGATT		7140
40					GTCTGACGCT		7200
					AAGGATCTTC		7260
					ATATĞAGTAA		7320
					GATCTGTCTA		7380
15					ACGGGAGGC		7440
45					GGCTCCAGAT		7500
					TGCAACTŤTA		7560
					TTCGCCAGTT		7620
					CTCGTCGTTŢ		7680
50					ATCCCCCATG\	N .	7740
50					TAAGTTGGCC	1	7800
					CATGCCATCC	1	7860
					ATAGTGTATG	\	7920
					ACATAGCAGA	\	7980
55					AAGGATCTTA	\	8040
55					TTCAGCATCT	1	8100
					CGCAAAAAAG	\	8160
					ATATTATTGA	\	8220
					TTAGAAAAAT	AAACAAATAG	8280
	GGGTTCCGCG	CACATTTCCC	CGAAAAGTGC	CACCTGACGT	CCBRAAG	\	8327

```
(2) INFORMATION FOR SEO ID NO:11:
            (i) SEQUENCE CHARACTERISTICS:
 5
              (A) \LENGTH: 8897 base pairs
              (B) TYPE: nucleic acid
              (C) STRANDEDNESS: single
              (D) TOPOLOGY: linear
10
            (ii) MOLECULE TYPE: cDNA
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:
      GGTACCAATT TAAATTGA\u00e7A TCTCCTTAGG TCTCGAGCAC CATGAAGTTG CCTGTTAGGC
15
      TGTTGGTGCT GATGTTCTĠĢ ATTCCTGCTT CCAGCAGTGA TGTTTTGATG ACCCAAATTC
      CAGTCTCCCT GCCTGTCAGT CTTGGAGATC AAGCGTCCAT CTCTTGCAGA TCTAGTCAGA
      TCATTGTACA TAATAATGGC \AACACCTATT TAGAATGGTA CCTGCAGAAA CCAGGCCAGT
      CTCCACAGCT CCTGATCTAC AAAGTTTCCA ACCGATTTTC TGGGGTCCCA GACAGGTTCA
      GCGGCAGTGG ATCAGGGACA GATTTCACAC TCAAGATCAG CAGAGTGGAG GCTGAGGATC
20
      TGGGAGTTTA TTACTGCTTT CAAGGTTCAC ATGTTCCATT CACGTTCGGC TCGGGGACAA
     AGTTGGAAAT AAAACGTAAG TCTCGAGTCT CTAGATAACC GGTCAATCGA TTGGAATTCT
     AAACTCTGAG GGGGTCGGAT GACGTGGCCA TTCTTTGCCT AAAGCATTGA GTTTACTGCA
      AGGTCAGAAA AGCATGCAAA GCCCCCAGAA TGGCTGCAAA GAGCTCCAAC AAAACAATTT
      AGAACTTTAT TAAGGAATAG GGGGAAGCTA GGAAGAAACT CAAAACATCA AGATTTTAAA
     TACGCTTCTT GGTCTCCTTG CTATAATTAT CTGGGATAAG CATGCTGTTT TCTGTCTGTC CCTAACATGC CCTTATCCGC AAACAAGCA CCCCAAGGGCA GAACTTTGTT ACTTAAACAC CATCCTGTTT GCTTCTTTCC TCAGGAAGTG TGGCTGCACC ATCTGTCTTC ATCTTCCCGC
      CATCTGATGA GCAGTTGAAA TCTGGAACAG CCTCTGTTGT GTGCCTGCTG AATAACTTCT
     ATCCCAGAGA GGCCAAAGTA CAGTGGAAGG TGGATAACGC CCTCCAATCG GGTAACTCCC
     AGGAGAGTGT CACAGAGCAG GAGAGCAAGG\ACAGCACCTA CAGCCTCAGC AGCACCCTGA
30
                                                                             1020
      CGCTGAGCAA AGCAGACTAC GAGAAACACA AAGTCTACGC CTGCGAAGTC ACCCATCAGG
      GCCTGAGCTC GCCCGTCACA AAGAGCTTCA ACAGGGGAGA GTGTTAGAGG GAGAAGTGCC
      CCCACCTGCT CCTCAGTTCC AGCCTGACCC COTCCCATCC TTTGGCCTCT GACCCTTTTT
      CCACAGGGGA CCTACCCCTA TTGCGGTCCT CCAGCTCATC TTTCACCTCA CCCCCCTCCT
35
      CCTCCTTGGC TTTAATTATG CTAATGTTGG AGGAGAATGA ATAAATAAAG TGAATCTTTG
                                                                             1320
      CACCTGTGGT TTCTCTCTTT CCTCATTTAA TAAT\ATTAT CTGTTGTTTT ACCAACTACT
      CAATTTCTCT TATAAGGGAC TAAATATGTA GTCATÇCTAA GGCACGTAAC CATTTATAAA
     AATCATCCTT CATTCTATTT TACCCTATCA TCCTCTGCAA GACAGTCCTC CCTCAAACCC
     ACAAGCCTTC TGTCCTCACA GTCCCCTGGG CCATGGTAGG AGAGACTTGC TTCCTTGTTT
40
     TCCCCTCCTC AGCAAGCCCT CATAGTCCTT TTTAAGGGTG ACAGGTCTTA CAGTCATATA
      TCCTTTGATT CAATTCCCTG AGAATCAACC AAAGCAAATT TTTCAAAAGA AGAAACCTGC
      TATAAAGAGA ATCATTCATT GCAACATGAT ATAAAATAAC AACACAATAA AAGCAATTAA
      ATAAACAAC AATAGGGAAA TGTTTAAGTT CATCATGGTA CTTAGACTTA ATGGAATGTC
     ATGCCTTATT TACATTTTTA AACAGGTACT GAGGGACTCC\TGTCTGCCAA GGGCCGTATT
45
      GAGTACTTTC CACAACCTAA TTTAATCCAC ACTATACTGT ÇAGATTAAAA ACATTCATTA
      AAATGTTGCA AAGGTTCTAT AAAGCTGAGA GACAAATATA TTCTATAACT CAGCAATCCC
      ACTTCTAGAT GACTGAGTGT CCCCACCCAC CAAAAAACTA TGCAAGAATG TTCAAAGCAG
      CTTTATTTAC AAAAGCCAAA AATTGGAAAT AGCCCGATTG TC\AACAATA GAATGAGTTA
                                                                             2100
      TTAAACTGTG GTATGTTTAT ACATTAGAAT ACCCAATGAG GAGAATTAAC AAGCTACAAC
                                                                             2160
50
      TATACCTACT CACACAGATG AATCTCATAA AAATAATGTT ACATAAGAGA AACTCAATGC
                                                                             2220
      AAAAGATATG TTCTGTATGT TTTCATCCAT ATAAAGTTCA AAACQAGGTA AAAATAAAGT
                                                                             2280
      TAGAAATTTG GATGGAAATT ACTCTTAGCT GGGGGTGGGC GAGTTAGTGC CTGGGAGAAG
      ACAAGAAGGG GCTTCTGGGG TCTTGGTAAT GTTCTGTTCC TCGTGTGGGG TTGTGCAGTT
      ATGATCTGTG CACTGTTCTG TATACACATT ATGCTTCAAA ATAACTTCAC ATAAAGAACA
55
      TCTTATACCC AGTTAATAGA TAGAAGAGGA ATAAGTAATA GGTCAAGACC AACGCAGCTG
      GTAAGTGGGG GCCTGGGATC AAATAGCTAC CTGCCTAATC CTGCCCWCTT GAGCCCTGAA
      TGAGTCTGCC TTCCAGGGCT CAAGGTGCTC AACAAACAA CAGGCCTGQT ATTTTCCTGG
      CATCTGTGCC CTGTTTGGCT AGCTAGGAGC ACACATACAT AGAAATTAAA TGAAACAGAC
                                                                             2700
      CTTCAGCAAG GGGACAGAGG ACAGAATTAA CCTTGCCCAG ACACTGGAAA CCCATGTATG
                                                                             2760
```

	AACACTCACA	TGTTTGGGAA	GGGGGAAGGG	CACATGTAAA	TGAGGACTCT	TCCTCATTCT	2820
			CCCTCTCAGC				2880
		1	TGAAGGGGTT				2940
	CAAATGACTG	ACAATCCCTT	TGTCCTGCTT	TGTTTTTCTT	TCCAGTCAGT	ACTGGGAAAG	3000
5			GAAACTACAT				3060
			TCTTTCAAAC				3120
	TAATGTCCCT	TCCAATGACA	TGAACTTGCT	CACTCATCCC	TGGGGGCCAA	ATTGAACAAT	3180
	CAAAGGCAGG	CATAATCCAG	TTATGAATTC	TTGCGGCCGC	TTGCTAGCTT	CACGTGTTGG	3240
	ATCCAACCGC	GGAAGGGCCC	TATTCTATAG	TGTCACCTAA	ATGCTAGAGC	TCGCTGATCA	3300
10	GCCTCGACTG	TGCCTTCTAG	TTGCCAGCCA	TCTGTTGTTT	GCCCTCCCC	CGTGCCTTCC	3360
	TTGACCCTGG	AAGGTGCCAG	TCCCACTGTC	CTTTCCTAAT	AAAATGAGGA	AATTGCATCG	3420
	CATTGTCTGA	GTAGGTGTCA	\TTCTATTCTG	GGGGGTGGGG	TGGGGCAGGA	CAGCAAGGGG	3480
	GAGGATTGGG	AAGACAATAG	CAGGCATGCT	GGGGATGCGG	TGGGCTCTAT	GGCTTCTGAG	3540
	GCGGAAAGAA	CCAGCTGGGG	CTÇTAGGGGG	TATCCCCACG	CGCCCTGTAG	CGGCGCATTA	3600
15	AGCGCGGCGG	GTGTGGTGGT	TACÇCGCAGC	GTGACCGCTA	CACTTGCCAG	CGCCCTAGCG	3660
			CCCTTCCTTT				3720
			CAATTAGTCA				3780
			CAGTTCCGCC				3840
	TTTATTTATG	CAGAGGCCGA	eeccecç4ce	GCCTCTGAGC	TATTCCAGAA	GTAGTGAGGA	3900
20	GGCTTTTTTG	GAGGCCTAGG	CTTTTGCAAA	AAGCTTGGAC	AGCTCAGGGC	TGCGATTTCG	3960
	CGCCAAACTT	${\tt GACGGCAATC}$	CTAGCGTGAA	GGCTGGTAGG	ATTTTATCCC	CGCTGCCATC	4020
	ATGGTTCGAC	CATTGAACTG	CATCGTCGCC	GTGTCCCAAA	ATATGGGGAT	TGGCAAGAAC	4080
			GCTCAGGAAC				4140
	ACCTCTTCAG	${\tt TGGAAGGTAA}$	ACAGAATCTG	ĠŢĠĀŢŦĀŢĠĠ	GTAGGAAAAC	CTGGTTCTCC	4200
25			TTTAAAGGAC	V /			4260
			TCATTTTCTT				4320
	ATTGAACAAC	CGGAATTGGC	AAGTAAAGTA	GAÇATGGTTT	GGATAGTCGG	AGGCAGTTCT	4380
			TCAACCAGGC				4440
••			${\tt GTTTTTCCCA}$				4500
30	CCAGAATACC	CAGGCGTCCT	CTCTGAGGTC	CAGGAGGAAA	AAGGCATCAA	GTATAAGTTT	4560
			CTAACAGGAA				4620
			ACCATGGGAC				4680
			ACATAATTGG	`			4740
2.5			AGTGTATAAT		\		4800
35			GGAACTGATG				4860
			GAAGAAATGC		\		4920
			AAAAAGAAGA		1		4980
			AGTCATGCTG		\		5040
40			AAAGCTGCAC		\		5100
40			CATAACAGTT				5160
			GCTATTAATA		`		5220
			AATAAGGAAT				5280
			TGTAGAGGTT				5340
15 .						TATTGCAGCT	5400
45					١	ATTTTTTCA	5460
			GTCCAAACTC				5520
						CAACTTGTTT	5580
			CAAATAAAGC				5640
50			TTGTGGTTTG				5700
50						GTTTCCTGTG	5760 5820
						AAAGTGTAAA	5820
			GAGCTAACTC			CGCGGGGAGA	5880 5940
			CTCTTCCGCT			1	6000
55						ATCCACAGAA	6060
55						CAGGAACCGT	6120
						GCATCACAAA	6180
						CCAGGCGTTT	6240
			GCGCTCTCCT				6300
	CCCCTGGAA	GCICCICGI	GCGCTCTCCT	O11CCGACCC	LUCCUCTIAC	COOMINGEIG	3300

```
TCCGCCTTTC 1CCCTTCGGG AAGCGTGGCG CTTTCTCAAT GCTCACGCTG TAGGTATCTC
                                                                             6360
      AGTTCGGTGT AGGTCGTTCG CTCCAAGCTG GGCTGTGTGC ACGAACCCCC CGTTCAGCCC
      GACCGCTGCG CCTTATCCGG TAACTATCGT CTTGAGTCCA ACCCGGTAAG ACACGACTTA
      TCGCCACTGG CAGQAGCCAC TGGTAACAGG ATTAGCAGAG CGAGGTATGT AGGCGGTGCT
 5
      ACAGAGTTCT TGAAQTGGTG GCCTAACTAC GGCTACACTA GAAGGACAGT ATTTGGTATC
      TGCGCTCTGC TGAAGCCAGT TACCTTCGGA AAAAGAGTTG GTAGCTCTTG ATCCGGCAAA
      CAAACCACCG CTGGTAGCGG TGGTTTTTTT GTTTGCAAGC AGCAGATTAC GCGCAGAAAA
      AAAGGATCTC AAGAAGAYCC TTTGATCTTT TCTACGGGGT CTGACGCTCA GTGGAACGAA
      AACTCACGTT AAGGGATTYT GGTCATGAGA TTATCAAAAA GGATCTTCAC CTAGATCCTT
10
      TTAAATTAAA AATGAAGTTY TAAATCAATC TAAAGTATAT ATGAGTAAAC TTGGTCTGAC
      AGTTACCAAT GCTTAATCAG TGAGGCACCT ATCTCAGCGA TCTGTCTATT TCGTTCATCC
      ATAGTTGCCT GACTCCCCGT CGTGTAGATA ACTACGATAC GGGAGGGCTT ACCATCTGGC
                                                                             7020
      CCCAGTGCTG CAATGATACC QCGAGACCCA CGCTCACCGG CTCCAGATTT ATCAGCAATA
      AACCAGCCAG CCGGAAGGGC CQAGCGCAGA AGTGGTCCTG CAACTTTATC CGCCTCCATC
      CAGTCTATTA ATTGTTGCCG GGAGCTAGA GTAAGTAGTT CGCCAGTTAA TAGTTTGCGC
15
      AACGTTGTTG CCATTGCTAC AGGCATCGTG GTGTCACGCT CGTCGTTTGG TATGGCTTCA
      TTCAGCTCCG GTTCCCAACG ATCAGGCGA GTTACATGAT CCCCCATGTT GTGCAAAAAA
                                                                             7320
      GCGGTTAGCT CCTTCGGTCC TCCGATCGTT GTCAGAAGTA AGTTGGCCGC AGTGTTATCA
                                                                             7380
      CTCATGGTTA TGGCAGCACT GCATAATTCT CTTACTGTCA TGCCATCCGT AAGATGCTTT
                                                                             7440
      TCTGTGACTG GTGAGTACTC AACCAAQTCA TTCTGAGAAT AGTGTATGCG GCGACCGAGT
20
      TGCTCTTGCC CGGCGTCAAT ACGGGATAAT ACCGCGCCAC ATAGCAGAAC TTTAAAAGTG
                                                                             7560
      CTCATCATTG GAAAACGTTC TTCGGGGCGA AAACTCTCAA GGATCTTACC GCTGTTGAGA
                                                                             7620
      TCCAGTTCGA TGTAACCCAC TCGTGCACCC AACTGATCTT CAGCATCTTT TACTTTCACC
      AGCGTTTCTG GGTGAGCAAA AACAGGAAGA CAAAATGCCG CAAAAAAGGG AATAAGGGCG
25
      ACACGGAAAT GTTGAATACT CATACTCTTC\CTTTTTCAAT ATTATTGAAG CATTTATCAG
                                                                             7800
      GGTTATTGTC TCATGAGCGG ATACATATTT &AATGTATTT AGAAAAATAA ACAAATAGGG
                                                                             7860
      GTTCCGCGCA CATTTCCCCG AAAAGTGCCA COTGACGTCG ACGGATCGGG AGATCTGCTA
GCCCGGGTGA CCTGAGGCGC GCCGGCTTCG AATAGCCAGA GTAACCTTTT TTTTTAATTT
                                                                            7920
      TATTTATTT TATTTTGAG ATGGAGTTTG GCGCCGATCT CCCGATCCCC TATGGTCGAC
30
      TCTCAGTACA ATCTGCTCTG ATGCCGCATA GTTAAGCCAG TATCTGCTCC CTGCTTGTGT
      GTTGGAGGTC GCTGAGTAGT GCGCGAGCAA AATTTAAGCT ACAACAAGGC AAGGCTTGAC
      CGACAATTGC ATGAAGAATC TGCTTAGGGT TAGGCCTTTTT GCGCTGCTTC GCGATGTACG
      GGCCAGATAT ACGCGTTGAC ATTGATTATT GACTAGTTAT TAATAGTAAT CAATTACGGG
      GTCATTAGTT CATAGCCCAT ATATGGAGTT CCGCGTTACA TAACTTACGG TAAATGGCCC
35
      GCCTGGCTGA CCGCCCAACG ACCCCCGCCC ATTGACGTCA ATAATGACGT ATGTTCCCAT
      AGTAACGCCA ATAGGGACTT TCCATTGACG TCAATGGGTG GACTATTTAC GGTAAACTGC
      CCACTTGGCA GTACATCAAG TGTATCATAT GCCAAGTACG CCCCCTATTG ACGTCAATGA
      CGGTAAATGG CCCGCCTGGC ATTATGCCCA GTACATGACC TTATGGGACT TTCCTACTTG
GCAGTACATC TACGTATTAG TCATCGCTAT TACCATGGTG ATGCGGTTTT GGCAGTACAT
40
      CAATGGGCGT GGATAGCGGT TTGACTCACG GGGATTTCCA AGTCTCCACC CCATTGACGT
      CAATGGGAGT TTGTTTTGGC ACCAAAATCA ACGGGACTTT CQAAAATGTC GTAACAACTC
      CGCCCCATTG ACGCAAATGG GCGGTAGGCG TGTACGGTGG GAGGTCTATA TAAGCAGAGC
                                                                             8820
      TCTCTGGCTA ACTAGAGAAC CCACTGCTTA CTGGCTTATC GAAATTAATA CGACTCACTA
                                                                             8880
      TAGGGAGACC CAAGCTT
45
               (2) INFORMATION FOR SEQ ID NO:12:
             (i) SEQUENCE CHARACTERISTICS:
              (A) LENGTH: 8321 base pairs
50
              (B) TYPE: nucleic acid
              (C) STRANDEDNESS: single
              (D) TOPOLOGY: linear
            (ii) MOLECULE TYPE: cDNA
55
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:
      GGTACCAATT TAAATTGATA TCTCCTTAGG TCTCGAGTCT CTAGATAACC GGTCAATCGA
                                                                              60
      TTGGAATTCT TGCGGCCGCT TGCTAGCCAC CATGGAGTTG TGGTTAAGCT TGGTCTTCCT
                                                                              120
```

		\					
	TGTCCTTGTT	TTAAAAGGTG	TCCAGTGTGA	AGTGCAACTG	GTGGAGTCTG	GGGGAGGCTT	180
	AGTGCAGCCT	GAGGGTCCC	TGCGACTTTC	CTGTGCTGCA	TCTGGATTCC	CGTTCAGTGA	240
		TATTGGGTTC					300
		GGTGATATAA					360
5		GCAAAGAACA					420
•		TACTOTGCAA					480
		GTCACGGTCT					540
		AAGAGCACCT					600
		CCGGTGACGG					660
10		GTCCTACAGT					720
10		TTGGGCACCC					780
		AAGAAAGTTG					840
		CGCTCCTGCC					900
		CGTCTGCCTC					960
15			1				1020
13		CTGGCTTTTT	1				
		ACACAAAGGG					1080
		CCCTGACCTA	1				1140
		CTCCTCCCAG	\				1200
20		AACTCACACA					1260
20		GCGGGACAGG	\				1320
		TCCGGAGCCA	\				1380
		TACCAACCTC	\				1440
		CCCGGGATGA	\				1500
0.5		CCAGCGACAT					1560
25		CGCCTCCCGT	17 %				1620
		AGAGCAGGTG					1680
		ACCACTACAC	1				1740
		GCCCCCGCTC					1800
• •		TACTTCCCGG					1860
30		ACTGTGATGG		1			1920
		CAGAGCGGGT		\			1980
	TGGGCCCCCT	AGGGTGGGC	TCAGCCAGGG	GCTGCCTCG	GCAGGGTGGG	GGATTTGCCA	2040
	GCGTGGCCCT	CCCTCCAGCA	GCACCTGCCC	TGGGCTGGGC	CACGGGAAGC	CCTAGGAGCC	2100
		GACACACAGC					2160
35	CTGTCCTCCC	GACCTCCATG	CCCACTCGGG	GGCATGCCTA	GTCCATGTGC	GTAGGGACAG	2220
	GCCCTCCCTC	ACCCATCTAC	CCCCACGGCA	CTAACCCCTG	GCTGCCCTGC	CCAGCCTCGC	2280
	ACCCGCATGG	GGACACAACC	GACTCCGGGG	ACATOCACTC	TCGGGCCCTG	TGGAGGGACT	2340
	GGTGCAGATG	CCCACACACA	CACTCAGCCC	AGACCGGTTC	AACAAACCCC	GCACTGAGGT	2400
	TGGCCGGCCA	CACGGCCACC	ACACACACAC	GTGCACGCCT	CACACACGGA	GCCTCACCCG	2460
40	GGCGAACTGC	ACAGCACCCA	GACCAGAGCA	AGGTCCTCGC	ACACGTGAAC	ACTCCTCGGA	2520
	CACAGGCCCC	CACGAGCCCC	ACGCGGCACC	TCAAGGCCCA	CGAGCCTCTC	GGCAGCTTCT	2580
	CCACATGCTG	ACCTGCTCAG	ACAAACCCAG	CCCTCCTCTC	ACAAGGGTGC	CCCTGCAGCC	2640
	GCCACACACA	CACAGGGGAT	CACACACCAC	GTCACGTCQC	TGGCCCTGGC	CCACTTCCCA	2700
	GTGCCGCCCT	TCCCTGCAGG	ACGGATCAGC	CTCGACTGTG	CCTTCTAGTT	GCCAGCCATC	2760
45	TGTTGTTTGC	CCCTCCCCCG	TGCCTTCCTT	GACCCTGGAA	GGTGCCACTC	CCACTGTCCT	2820
	TTCCTAATAA	AATGAGGAAA	TTGCATCGCA	TTGTCTGAGT	AGGTGTCATT	CTATTCTGGG	2880
					1	GGCATGCTGG	2940
	GGATGCGGTG	GGCTCTATGG	CTTCTGAGGC	GGAAAGAACC	AGCTGGGGCT	CTAGGGGGTA	3000
	TCCCCACGCG	CCCTGTAGCG	GCGCATTAAG	CGCGGCGGGT	GTGGTGA	CGCGCAGCGT	3060
50		CTTGCCAGCG					3120
		GCCGGGCCTC					3180
					1	GTTCCGCCCA	3240
		CATGGCTGAC			``		3300
						TTTGCAAAAA	3360
55		CTCAGGGCTG					3420
		TTTATCCCCG					3480
						TCAGGAACGA	3540
		TTCCAAAGAA					3600
		AGGAAAACCT					3660
					1		

	\						
	ATATTAA	GTTCTCAGTA	GAGAACTCAA	AGAACCACCA	CGAGGAGCTC	ATTTTCTTGC	3720
	CAAAAGTTTG	GATGATGCCT	TAAGACTTAT	TGAACAACCG	GAATTGGCAA	GTAAAGTAGA	3780
	CATGGTTTOG	ATAGTCGGAG	GCAGTTCTGT	TTACCAGGAA	GCCATGAATC	AACCAGGCCA	3840
	CCTTAGACTA	TTTGTGACAA	GGATCATGCA	GGAATTTGAA	AGTGACACGT	TTTTCCCAGA	3900
5	AATTGATTTG	GGGAAATATA					3960
		GCATCAAGT					4020
		TACTCTGCTC					4080
		TAGATCTCTT					4140
		CAGAGATTTA					4200
10		TGATTCTAAT					4260
• •		GGTGGAATGC					4320
		ATGAGGCTAC					4320
		ACCCCAAGGA					4440
		GAACTCTTGC					4500
15		AAATTATGGA					4560
13		TACTGTTTTT	1				
			\				4620
		AATTGTGTAC	\				4680
		GTGCCTTGAC					4740
20		AAAAACCTCC					4800
20		AACTTGTTTA					4860
		AATAAAGCAT	\ \				4920
		TATCATGTCT	\				4980
		GCCCACCCCA	\				5040
25		AATTTCACAA	\				5100
25		AATGTATCTT	\ \ \	i e			5160
		TCATAGCTGT					5220
		GGAAGCATAA	صيانا	\			5280
		TTGCGCTCAC		\			5340
20		GGCCAACGCG		`			5400
30		GACTCGCTGC					5460
	AAAGGCGGTA	ATACGGTTAT	CCACAGAATC	AGGGATAAC	GCAGGAAAGA	ACATGTGAGC	5520
		CAAAAGGCCA		\			5580
	GCTCCGCCCC	CCTGACGAGC	ATCACAAAAA	TCGAGGCTCA	AGTCAGAGGT	GGCGAAACCC	5640
	GACAGGACTA	TAAAGATACC	AGGCGTTTCC	CCCTGGAAGC	TCCCTCGTGC	GCTCTCCTGT	5700
35	TCCGACCCTG	CCGCTTACCG	GATACCTGTC	CGCCTTTCTC	CCTTCGGGAA	GCGTGGCGCT	5760
	TTCTCAATGC	TCACGCTGTA	GGTATCTCAG	TTCGGTCTAG	GTCGTTCGCT	CCAAGCTGGG	5820
	CTGTGTGCAC	GAACCCCCCG	TTCAGCCCGA	CCGCTGCGCC	TTATCCGGTA	ACTATCGTCT	5880
	TGAGTCCAAC	CCGGTAAGAC	ACGACTTATC	GCCACTGGGA	GCAGCCACTG	GTAACAGGAT	5940
	TAGCAGAGCG	AGGTATGTAG	GCGGTGCTAC	AGAGTTCTTC	AAGTGGTGGC	CTAACTACGG	6000
40	CTACACTAGA	AGGACAGTAT	TTGGTATCTG	CGCTCTGCTG\	\AAGCCAGTTA	CCTTCGGAAA	6060
	AAGAGTTGGT	AGCTCTTGAT	CCGGCAAACA	AACCACCGCT	GTAGCGGTG	GTTTTTTTGT	6120
	TTGCAAGCAG	CAGATTACGC	GCAGAAAAA	AGGATCTCAA	GAAGATCCTT	TGATCTTTTC	6180
	TACGGGGTCT	GACGCTCAGT	GGAACGAAAA	CTCACGTTAA	GGÇATTTTGG	TCATGAGATT	6240
	ATCAAAAAGG	ATCTTCACCT	AGATCCTTTT	AAATTAAAA	TGAAGTTTTA	AATCAATCTA	6300
45	AAGTATATAT	GAGTAAACTT	GGTCTGACAG	TTACCAATGC	TTAATCAGTG	AGGCACCTAT	6360
	CTCAGCGATC	TGTCTATTTC	GTTCATCCAT	AGTTGCCTGA	стссфсстсс	TGTAGATAAC	6420
	TACGATACGG	GAGGGCTTAC	CATCTGGCCC	CAGTGCTGCA	ATGATACCGC	GAGACCCACG	6480
	CTCACCGGCT	CCAGATTTAT	CAGCAATAAA	CCAGCCAGCC	GGAAGGGCCG	AGCGCAGAAG	6540
		ACTTTATCCG					6600
50	AAGTAGTTCG	CCAGTTAATA	GTTTGCGCAA	CGTTGTTGCC	ATTGCTACAG	GCATCGTGGT	6660
	GTCACGCTCG	TCGTTTGGTA	TGGCTTCATT	CAGCTCCGGT	TCCCAACGAT	CAAGGCGAGT	6720
		CCCATGTTGT			1		6780
		TTGGCCGCAG			\		6840
		CCATCCGTAA				t	6900
55	CTGAGAATAG	TGTATGCGGC	GACCGAGTTG	CTCTTGCCCG	GCGTCAATAC	GGATAATAC	. 6960
		AGCAGAACTT				\	7020
		ATCTTACCGC					7080
		GCATCTTTTA					7140
		AAAAAGGGAA				\	7200
			-			\	

30

```
TTTTCAATAT TATTGAAGCA TTTATCAGGG TTATTGTCTC ATGAGCGGAT ACATATTTGA
                                                                              7260
      ATGTATTTAG AAAAATAAAC AAATAGGGGT TCCGCGCACA TTTCCCCGAA AAGTGCCACC
                                                                              7320
      TGACGTCGAC\GGATCGGGAG ATCTGCTAGG TGACCTGAGG CGCGCCGGCT TCGAATAGCC
                                                                              7380
      AGAGTAACCT TTTTTTTAA TTTTATTTTA TTTTATTTTT GAGATGGAGT TTGGCGCCGA
 5
      TCTCCCGATC CCTATGGTC GACTCTCAGT ACAATCTGCT CTGATGCCGC ATAGTTAAGC
                                                                              7500
      CAGTATCTGC TCCCTGCTTG TGTGTTGGAG GTCGCTGAGT AGTGCGCGAG CAAAATTTAA
                                                                              7560
      GCTACAACAA GGCAAGGCTT GACCGACAAT TGCATGAAGA ATCTGCTTAG GGTTAGGCGT
                                                                              7620
      TTTGCGCTGC TTCQCGATGT ACGGGCCAGA TATACGCGTT GACATTGATT ATTGACTAGT
      TATTAATAGT AATCAATTAC GGGGTCATTA GTTCATAGCC CATATATGGA GTTCCGCGTT
10
      ACATAACTTA CGGTAAATGG CCCGCCTGGC TGACCGCCCA ACGACCCCCG CCCATTGACG
      TCAATAATGA CGTATGATCC CATAGTAACG CCAATAGGGA CTTTCCATTG ACGTCAATGG
      GTGGACTATT TACGGTAAAC TGCCCACTTG GCAGTACATC AAGTGTATCA TATGCCAAGT
      ACGCCCCTA TTGACGTCAA TGACGGTAAA TGGCCCGCCT GGCATTATGC CCAGTACATG
                                                                              7980
      ACCTTATGGG ACTTTCCTAC TTGGCAGTAC ATCTACGTAT TAGTCATCGC TATTACCATG GTGATGCGGT TTTGGCAGTA CATCAATGGG CGTGGATAGC GGTTTGACTC ACGGGGATTT
                                                                              8040
15
                                                                              8100
      CCAAGTCTCC ACCCCATTGA GTCAATGGG AGTTTGTTTT GGCACCAAAA TCAACGGGAC
                                                                              8160
      TTTCCAAAAT GTCGTAACAA CTCCGCCCCA TTGACGCAAA TGGGCGGTAG GCGTGTACGG
                                                                              8220
      TGGGAGGTCT ATATAAGCAG AGTCTCTGG CTAACTAGAG AACCCACTGC TTACTGGCTT
                                                                              8280
      ATCGAAATTA ATACGACTCA CTATAGGGAG ACCCAAGCTT G
                                                                              8321
20
                (2) INFORMATION FOR SEQ ID NO:13:
            (i) SEQUENCE CHARACTERISTICS:
              (A) LENGTH: 8897 base pairs
25
               (B) TYPE: nucleic acid
              (C) STRANDEDNESS: single
              (D) TOPOLOGY: linear
```

(ii) MOLECULE TYPE: cDNA

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

	GACGGATCGG	GAGATCTGCT	AGCCCGGGTG	ACCT@AGGCG	CGCCGGCTTC	GAATAGCCAG	60
	AGTAACCTTT	${\tt TTTTTTTAATT}$	TTATTTTATT	TTATTTTTGA	GATGGAGTTT	GGCGCCGATC	120
35	TCCCGATCCC	${\tt CTATGGTCGA}$	CTCTCAGTAC	AATCTGCTCT	GATGCCGCAT	AGTTAAGCCA	180
	GTATCTGCTC	CCTGCTTGTG	TGTTGGAGGT	CGCTGAGTAG	TGCGCGAGCA	AAATTTAAGC	240
	TACAACAAGG	CAAGGCTTGA	CCGACAATTG	CATGAAGAAT	CTGCTTAGGG	TTAGGCGTTT	300
	TGCGCTGCTT	CGCGATGTAC	GGGCCAGATA	TACGCGTTOA	CATTGATTAT	TGACTAGTTA	360
	TTAATAGTAA	${\tt TCAATTACGG}$	GGTCATTAGT	TCATAGCCCA	TATATGGAGT	TCCGCGTTAC	420
40	ATAACTTACG	${\tt GTAAATGGCC}$	CGCCTGGCTG	ACCGCCCAAC	GACCCCCGCC	CATTGACGTC	480
	AATAATGACG	TATGTTCCCA	TAGTAACGCC	AATAGGGACT	TTCCATTGAC	GTCAATGGGT	540
	GGACTATTTA	CGGTAAACTG	CCCACTTGGC	AGTACATCAA	GTATCATA	TGCCAAGTAC	600
	GCCCCCTATT	GACGTCAATG	ACGGTAAATG	GCCCGCCTGG	CATTATGCCC	AGTACATGAC	660
	CTTATGGGAC	TTTCCTACTT	GGCAGTACAT	CTACGTATTA	GTCATCGCTA	TTACCATGGT	720
45	GATGCGGTTT	TGGCAGTACA	TCAATGGGCG	TGGATAGCGG	TTTGACTCAC	GGGGATTTCC	780
	AAGTCTCCAC	CCCATTGACG	TCAATGGGAG	TTTGTTTTGG	CACCAAAATC	AACGGGACTT	840
	TCCAAAATGT	CGTAACAACT	CCGCCCCATT	GACGCAAATG	GGCGGTAGGC	GTGTACGGTG	900
	GGAGGTCTAT	ATAAGCAGAG	CTCTCTGGCT	AACTAGAGAA	CCCACTGCTT	ACTGGCTTAT	960
	CGAAATTAAT	ACGACTCACT	ATAGGGAGAC	CCAAGCTTGG	TACCAATTTA	AATTGATATC	1020
50	TCCTTAGGTC	TCGAGCACCA	TGAAGTTGCC	TGTTAGGCTG	TTGGTGTTGA	TGTTCTGGAT	1080
	TCCTGCTTCC	AGCAGTGATG	TTGTCATGAC	CCAAACCCCA	CTGTCCAGTC	CTGTCACGCT	1140
	TGGACAACCT	GCGTCCATCT	CTTGCAGATC	TAGTCAGATC	ATTGTACATA	ATAATGGCAA	1200
	CACCTATCTG	GAATGGTACC	AGCAGAGACC	AGGGCAGTCT	CCACGGCTQC	TGATCTACAA	1260
	AGTTTCCAAC	CGATTTTCTG	GGGTCCCAGA	CAGGTTCAGC	GGCAGTGGAG	CTGGGACAGA	1320
55	TTTCACACTC	AAGATCAGCA	GAGTGGAGGC	TGAGGATGTG	GGAGTTTACT\	ACTGCTTCCA	1380
	GGGTTCACAT	GTTCCATTCA	CGTTCGGCCA	AGGGACAAAG	TTGGAAATCA	PACGTAAGTC	1440
	TCGAGTCTCT	AGATAACCGG	TCAATCGATT	GGAATTCTAA	ACTCTGAGGG	dGTCGGATGA	1500
	CGTGGCCATT	CTTTGCCTAA	AGCATTGAGT	TTACTGCAAG	GTCAGAAAAG	CATGCAAAGC	1560
	CCTCAGAATG	GCTGCAAAGA	GCTCCAACAA	AACAATTTAG	AACTTTATTA	ACCAATAGGG	1620

CARCCATICG CAGATICTICT ACCCCCCTTC ACCTRACGC CCCCCCTTC CAATACCCAC

